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Sequence Listing could not be accepted.

If you need help call the Patent Electronic Business Center at (866)
217-9197 (toll free).

Reviewer: Anne Corrigan

Timestamp: [year=2008; month=5; day=1; hr=18; min=58; sec=33; ms=853;]

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Reviewer Comments:

SUBSTITUTE SEQUENCE LISTING

Please remove "SUBSTITUTE" above; just use "SEQUENCE LISTING"

<210> 12

<211> 16

<212> PRT

<213> Artificial Sequence

<220>

<223> Synthetic Peptide

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<223> Xaa is any amino acid

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The above <220>-<223> section describing Xaa at location 17 is incorrect: only 16 amino acids are in this sequence.

<210> 13
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<212> DNA
<213> Synthetic Construct

<220>

<221> CDS

<222> (1)..(1833)

<400> 13

The above <213> response is invalid, per 1.823 of Sequence Rules. The only valid responses are: the Genus species of the organism, "Artificial Sequence," or "Unknown." "Artificial Sequence" and "Unknown" require explanation in the <220>-<223> section: please give the source of the genetic material. Same error in Sequence 14.

Application No: 10540247 Version No: 2.0

Input Set:

Output Set:

Started: 2008-04-21 16:37:52.866
Finished: 2008-04-21 16:37:57.988
Elapsed: 0 hr(s) 0 min(s) 5 sec(s) 122 ms
Total Warnings: 14
Total Errors: 0
No. of SeqIDs Defined: 14
Actual SeqID Count: 14

| Error code | Error Description |
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| W 402 | Undefined organism found in <213> in SEQ ID (13) |
| W 402 | Undefined organism found in <213> in SEQ ID (14) |

SUBSTITUTE SEQUENCE LISTING

<110> DE WAARD, Michel
 DUPUIS, Alain
 GRUNWALD, Didier
 SANDOZ, Guillaume

<120> CHIMERIC PROTEIN FOR THE SCREENING OF AGONISTS AND ANTAGONISTS OF
 CELL SIGNALLING PATHWAYS THAT ARE DEPENDENT ON G-PROTEIN-COUPLED
 RECEPTORS

<130> 273623US0XPCT

<140> 10540247

<141> 2006-06-05

<150> PCT/FR03/003860

<151> 2003-12-22

<150> FR 02/16576

<151> 2002-12-23

<160> 14

<170> PatentIn version 3.3

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<211> 51

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| aattgcatcg cattgtctga gtaggtgtca ttctattctg gggggtgggg tggggcagga | 2100 |
| cagcaagggg gaggattggg aagacaatag caggcatgct ggggatgcgg tgggctctat | 2160 |

| | |
|--|------|
| ggcttctgag gcggaagaa ccagctgggg ctctagggg tatccccacg cgccctgtag | 2220 |
| cggcgcatta agcgcggcgg gtgtgggtgt tacgcgcagc gtgaccgcta cacttgccag | 2280 |
| cgccctagcg cccgctcctt tcgctttctt cccttccttt ctcgccacgt tcgcgggctt | 2340 |
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| cctcgacccc aaaaaacttg attagggtga tggttcacgt agtgggccat cgccctgata | 2460 |
| gacggttttt cgcccttga cgttgagtc cacgttcttt aatagtggac tcttgttcca | 2520 |
| aactggaaca acactcaacc ctatctcggc ctattctttt gatttataag ggattttggg | 2580 |
| gatttcggcc tattgggttaa aaaatgagct gatttaacaa aaatttaacg cgaattaatt | 2640 |
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| agcaggcaga agtatgcaaa gcatgcatct caattagtca gcaaccatag tcccgccct | 2820 |
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| atccattttc ggatctgac aagagacagg atgaggatcg ttctgcatga ttgaacaaga | 3060 |
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| attcatcgac tgtggccggc tgggtgtggc ggaccgctat caggacatag cgttggctac | 3720 |
| ccgtgatatt gctgaagagc ttggcggcga atgggctgac cgcttcctcg tgctttacgg | 3780 |
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| agcgggactc tgggggttca aatgaccgac caagcgacgc ccaacctgcc atcacgagat | 3900 |

| | |
|--|------|
| ttcgattcca cgcgcgctt ctatgaaagg ttgggcttcg gaatcgtttt ccgggacgcc | 3960 |
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| tttattgcag cttataatgg ttacaaataa agcaatagca tcacaaattt cacaaataaa | 4080 |
| gcattttttt cactgcattc tagttgtggt ttgtccaaac tcatcaatgt atcttatcat | 4140 |
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| gtgtgaaatt gttatccgct cacaattcca cacaacatac gagccggaag cataaagtgt | 4260 |
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| aaaa | |